

Figure 1

>Alignment of Atlno 80 sequence and public sequence, At3g57300, showing splicing difference

Query: claimed sequence

Sbjct: gi|18410689|ref|NM\_115590.1| (AGI:At3g57300)

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Query: 3961 acaaaaagatg 3970  
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Sbjct: 3961 acaaaaagatg 3970

Score = 1001 bits (505), Expect = 0.0  
Identities = 522/528 (98%), Gaps = 6/528 (1%)  
Strand = Plus / Plus

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Sbjct: 3970 gagacggtggaagagaaaattttgcacagggcaagtcagaaaaatacagttcaacagctt 4029

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